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QM protein - protein search, using sw model

Run on: January 16, 2003, 16:44:27 - Search time 3.0712 seconds
                                        (without alignments)
                                        32.366 Million Cell updates/sec

Title: US-09-856-070-18
Perfect score: 24
Sequence: 1 KERLM S

Scoring table: BLOSUM62
Gap-op 10.0, Gap-ext 0.5

Searched: 120991 seqs. 19878514 residues

Total number of hits satisfying chosen parameters: 120991

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
                 Maximum Match 100%
                 Listing first 45 summaries

Database :   Published_Applications_AA:*
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    2: /cgn2_6/ptodata/2/pubpaa/PCT_NEW_PUB.pep:*
    3: /cgn2_6/ptodata/2/pubpaa/US06_NEW_PUB.pep:*
    4: /cgn2_6/ptodata/2/pubpaa/US06_PUBCOMB.pep:*
    5: /cgn2_6/ptodata/2/pubpaa/US07_NEW_PUB.pep:*
    6: /cgn2_6/ptodata/2/pubpaa/US07_PUBCOMB.pep:*
    7: /cgn2_6/ptodata/2/pubpaa/PCTUS_PUBCOMB.pep:*
    8: /cgn2_6/ptodata/2/pubpaa/US08_PUBCOMB.pep:*
    9: /cgn2_6/ptodata/2/pubpaa/US09_NEW_PUB.pep:*
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   11: /cgn2_6/ptodata/2/pubpaa/US10_NEW_PUB.pep:*
   12: /cgn2_6/ptodata/2/pubpaa/US10_PUBCOMB.pep:*
   13: /cgn2_6/ptodata/2/pubpaa/US60_NEW_PUB.pep:*
   14: /cgn2_6/ptodata/2/pubpaa/US60_PUBCOMB.pep:*

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	24	100.0	29	10	US-09-864-761-37381	Sequence 37381, A
2	24	100.0	63	10	US-09-385-903-59	Sequence 59, Appl
3	24	100.0	68	10	US-09-704-869-652	Sequence 652, App
4	24	100.0	164	10	US-09-815-242-11411	Sequence 11411, A
5	24	100.0	164	10	US-09-815-242-11573	Sequence 11573, A
6	24	100.0	171	10	US-09-888-243-13	Sequence 13, Appl
7	24	100.0	172	10	US-09-888-243-24	Sequence 24, Appl
8	24	100.0	172	10	US-09-888-243-27	Sequence 27, Appl
9	24	100.0	434	9	US-10-060-425-15	Sequence 15, Appl
10	24	100.0	434	10	US-09-827-040-7	Sequence 7, Appli
11	24	100.0	451	10	US-09-888-243-28	Sequence 28, Appl
12	24	100.0	476	9	US-10-060-425-4	Sequence 4, Appli
13	24	100.0	476	9	US-10-060-425-12	Sequence 12, Appl
14	24	100.0	476	9	US-10-060-425-14	Sequence 14, Appl
15	24	100.0	476	10	US-09-827-040-2	Sequence 2, Appli
16	24	100.0	476	10	US-09-827-040-3	Sequence 3, Appli
17	24	100.0	476	10	US-09-827-040-6	Sequence 6, Appli
18	24	100.0	482	10	US-09-804-551B-32	Sequence 32, Appl
19	24	100.0	491	10	US-09-925-300-1715	Sequence 1715, Ap

20	24	100.0	625	10	US-09-925-300-896	Sequence 896, App
21	24	100.0	1150	10	US-09-870-122-3	Sequence 3, Appli
22	24	100.0	1154	10	US-09-870-122-1	Sequence 1, Appli
23	24	100.0	1167	10	US-09-870-122-2	Sequence 2, Appli
24	24	100.0	1181	10	US-09-870-122-23	Sequence 23, Appl
25	24	100.0	1215	10	US-09-817-913-11	Sequence 11, Appl
26	24	100.0	1215	10	US-09-976-280A-4	Sequence 4, Appli
27	24	100.0	1215	10	US-09-976-280A-22	Sequence 22, Appl
28	24	100.0	1215	10	US-09-817-538-11	Sequence 11, Appl
29	24	91.7	47	10	US-09-864-761-42087	Sequence 42087, A
30	22	91.7	203	9	US-09-350-874-12	Sequence 12, Appl
31	22	91.7	206	10	US-09-965-528-15	Sequence 15, Appl
32	22	91.7	216	10	US-09-350-874-2	Sequence 2, Appli
33	22	91.7	216	10	US-09-350-874-6	Sequence 6, Appli
34	22	91.7	227	10	US-09-350-874-8	Sequence 8, Appli
35	22	91.7	227	10	US-09-350-874-10	Sequence 10, Appl
36	22	91.7	245	10	US-09-350-874-4	Sequence 4, Appli
37	21	87.5	35	10	US-09-864-761-42517	Sequence 42517, A
38	21	87.5	55	10	US-09-864-761-194-17	Sequence 17, Appl
39	21	87.5	67	10	US-09-996-194-17	Sequence 45598, A
40	21	87.5	68	10	US-09-864-761-45598	Sequence 34946, A
41	21	87.5	86	10	US-09-864-761-34946	Sequence 48121, A
42	21	87.5	86	10	US-09-864-761-48121	Sequence 46, Appl
43	21	87.5	87	10	US-09-728-721-46	Sequence 24, Appl
44	21	87.5	92	10	US-09-916-740-24	Sequence 46056, A
45	21	87.5	95	10	US-09-864-761-46056	Sequence 1/127, A

ALIGNMENTS

RESULT 1
US-09-864-761-37381

  Sequence 37381, Application US09864761
  Patent No. US2003004876341
  GENERAL INFORMATION:
    APPLICANT: Penn, Sharron G.
    APPLICANT: Funk, David K.
    APPLICANT: Hauzel, David K.
    APPLICANT: Chen, Wensheng
    TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FO
    TITLE OF INVENTION: GENE EXPRESSION ANALYSIS IN MICROARRAY
    FILE REFERENCE: Acomica-X-1
    CURRENT APPLICATION NUMBER: US-09/864,761
    CURRENT FILING DATE: 2001-05-23
    PRIOR APPLICATION NUMBER: US 60/180,312
    PRIOR FILING DATE: 2000-02-04
    PRIOR APPLICATION NUMBER: US 60/207,456
    PRIOR FILING DATE: 2000-05-26
    PRIOR APPLICATION NUMBER: US 09/632,366
    PRIOR FILING DATE: 2000-08-03
    PRIOR APPLICATION NUMBER: GB 24261.6
    PRIOR FILING DATE: 2000-10-04
    PRIOR APPLICATION NUMBER: US 60/236,359
    PRIOR FILING DATE: 2000-09-27
    PRIOR APPLICATION NUMBER: PCT/US01/00666
    PRIOR FILING DATE: 2001-01-30
    PRIOR APPLICATION NUMBER: PCT/US01/00667
    PRIOR FILING DATE: 2001-01-30
    PRIOR APPLICATION NUMBER: PCT/US01/00664
    PRIOR FILING DATE: 2001-01-30
    PRIOR APPLICATION NUMBER: PCT/US01/00669
    PRIOR FILING DATE: 2001-01-30
    PRIOR APPLICATION NUMBER: PCT/US01/00665
    PRIOR FILING DATE: 2001-01-30
    PRIOR APPLICATION NUMBER: PCT/US01/00668
    PRIOR FILING DATE: 2001-01-30
    PRIOR APPLICATION NUMBER: PCT/US01/00663
    PRIOR FILING DATE: 2001-01-30
    PRIOR APPLICATION NUMBER: PCT/US01/00662
    PRIOR FILING DATE: 2001-01-30
    PRIOR APPLICATION NUMBER: PCT/US01/00661
    PRIOR FILING DATE: 2001-01-30

; PRIOR APPLICATION NUMBER: PCT/US01/00670
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: US 60/234,687
; PRIOR FILING DATE: 2000-09-21
; PRIOR APPLICATION NUMBER: US 09/608,408
; PRIOR FILING DATE: 2000-06-30
; PRIOR APPLICATION NUMBER: US 09/774,204
; PRIOR FILING DATE: 2001-01-29
; NUMBER OF SEQ ID NOS: 49117
; SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
; SEQ ID NO 37481
; LENGTH: 29
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; OTHER INFORMATION: MAP IG A1031116.1
; OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL - 1.8
; OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL - 2.1
; OTHER INFORMATION: EXPRESSED IN HH1100, SIGNAL - 1.8
; OTHER INFORMATION: EXPRESSED IN BT474, SIGNAL - 1.9
; OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL - 1.9
; OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL - 2.5
; OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL - 2
; OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL - 1.6
; OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL - 2.2
; OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL - 1.9
; OTHER INFORMATION: BSI_HUMAN H11: AW452877.1, EVALUE 9.00e-01
US-09-864-761-37481

Query Match   100.0%; Score 24; DB 10; Length 29;
Best Local Similarity 100.0%; Pred. No. 9.3;
Matches   5; Conservative   0; Mismatches   0; Indels   0; Gaps   0;

Qy   1 KEELM 5
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Db   8 KEELM 12

RESULT 2
US-09-989-903-59, Application US/09989903
; Sequence 59, Application US/09989903
; Patent No. US20020146804A1
; GENERAL INFORMATION:
; APPLICANT: Alnemri, Emad S.
; APPLICANT: Fernandez Alnemri, Teresa
; TITLE OF INVENTION: CASPASE-14, AN APOPTOTIC PROTEASE, NUCLEIC ACID ENCODING AND METHODS OF USE.
; FILE REFERENCE: 480140.434D1
; CURRENT APPLICATION NUMBER: US/09/989,903
; CURRENT FILING DATE: 2002-04-11
; NUMBER OF SEQ ID NOS: 78
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 59
; LENGTH: 53
; TYPE: PRT
; ORGANISM: Mus musculus
US-09-989-903-59

Query Match   100.0%; Score 24; DB 10; Length 53;
Best Local Similarity 100.0%; Pred. No. 18;
Matches   5; Conservative   0; Mismatches   0; Indels   0; Gaps   0;

Qy   1 KEELM 5
        |||||
Db   5 KEELM 9

RESULT 4
US-09-764-869-652
; Sequence 652, Application US/09764869
; Patent No. US2002006152IA1
; GENERAL INFORMATION:
; APPLICANT: Rosen et al.

; TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
; FILE REFERENCE: PC007
; CURRENT APPLICATION NUMBER: US/09/764,869
; CURRENT FILING DATE: 2001-01-17
; Prior application data removed   refer to PALM or file wrapper
; NUMBER OF SEQ ID NOS: 2442
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 652
; LENGTH: 68
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (15)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
US-09-764-869-652

Query Match   100.0%; Score 24; DB 10; Length 68;
Best Local Similarity 100.0%; Pred. No. 23;
Matches   5; Conservative   0; Mismatches   0; Indels   0; Gaps   0;

Qy   1 KEELM 5
        |||||
Db 27 KEELM 31

RESULT 4
US-09-815-242-11411
; Sequence 11411, Application US/09815242
; Patent No. US20020015691A1
; GENERAL INFORMATION:
; APPLICANT: Haselbeck, Robert
; APPLICANT: Ohlsen, Karl L.
; APPLICANT: Zyskind, Judith W.
; APPLICANT: Wall, Daniel
; APPLICANT: Trawick, John D.
; APPLICANT: Carr, Grant J.
; APPLICANT: Yamamoto, Robert T.
; APPLICANT: Xu, H. Howard
; TITLE OF INVENTION: Identification of Essential Genes in Prokaryotes
; FILE REFERENCE: ELITRA.011A
; CURRENT APPLICATION NUMBER: US/09/815,242
; CURRENT FILING DATE: 2001-03-21
; PRIOR APPLICATION NUMBER: 60/191,078
; PRIOR FILING DATE: 2000-03-21
; PRIOR APPLICATION NUMBER: 60/206,848
; PRIOR FILING DATE: 2000-05-23
; PRIOR APPLICATION NUMBER: 60/207,727
; PRIOR FILING DATE: 2000-05-26
; PRIOR APPLICATION NUMBER: 60/242,578
; PRIOR FILING DATE: 2000-10-23
; PRIOR APPLICATION NUMBER: 60/253,625
; PRIOR FILING DATE: 2000-11-27
; PRIOR APPLICATION NUMBER: 60/257,931
; PRIOR FILING DATE: 2000-12-22
; PRIOR APPLICATION NUMBER: 60/269,308
; PRIOR FILING DATE: 2001-02-16
; NUMBER OF SEQ ID NOS: 14110
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 11411
; LENGTH: 164
; TYPE: PRT
; ORGANISM: Helicobacter pylori
US-09-815-242-11411

Query Match   100.0%; Score 24; DB 10; Length 164;
Best Local Similarity 100.0%; Pred. No. 58;
Matches   5; Conservative   0; Mismatches   0; Indels   0; Gaps   0;

Qy   1 KEELM 5
        |||||
Db 132 KEELM 136

RESULT 5
US-09-815-242-11573
; Sequence 11573, Application US/09815242
; Patent No. US20020061569A1
; GENERAL INFORMATION:
; APPLICANT: Haselbeck, Robert
; APPLICANT: Ohlsen, Kari L.
; APPLICANT: Zyskind, Judith W.
; APPLICANT: Wall, Daniel
; APPLICANT: Trawick, John D.
; APPLICANT: Carr, Grant J.
; APPLICANT: Yamamoto, Robert T.
; APPLICANT: Xu, H. Howard
; TITLE OF INVENTION: Identification of Essential Genes in
; TITLE OF INVENTION: Prokaryotes
; FILE REFERENCE: ELITRA.011A
; CURRENT APPLICATION NUMBER: US/09,815,242
; CURRENT FILING DATE: 2001-03-21
; PRIOR APPLICATION NUMBER: 60/191,078
; PRIOR FILING DATE: 2000-03-21
; PRIOR APPLICATION NUMBER: 60/206,848
; PRIOR FILING DATE: 2000-05-23
; PRIOR APPLICATION NUMBER: 60/207,727
; PRIOR FILING DATE: 2000-05-26
; PRIOR APPLICATION NUMBER: 60/242,578
; PRIOR FILING DATE: 2000-10-23
; PRIOR APPLICATION NUMBER: 60/253,625
; PRIOR FILING DATE: 2000-11-27
; PRIOR APPLICATION NUMBER: 60/257,991
; PRIOR FILING DATE: 2000-12-22
; PRIOR APPLICATION NUMBER: 60/259,408
; PRIOR FILING DATE: 2001-01-02
; NUMBER OF SEQ ID NOS: 14110
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 11573
; LENGTH: 164
; TYPE: PRT
; ORGANISM: Helicobacter pylori
US-09-815-242-11573

Qry Match.   100.0%, Score 24; DB 10, Length 164,
Best Local Similarity 100.0%, Prd. No. 58;
Matches   5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy   1 KEELM 5
         |||||
Db   132 KEELM 136

RESULT 6
US-09-888-243-13
; Sequence 13, Application US/09888243
; Patent No. US20020136714A1
; GENERAL INFORMATION:
; APPLICANT: Horvitz, H. Robert
; APPLICANT: Yuan, Junying
; APPLICANT: Shaham, Shai
; TITLE OF INVENTION: Relatedness of Human Interleukin-1beta
; TITLE OF INVENTION: Convertase Gene to a C. Elegans Cell Death Gene, Inhibitory
; TITLE OF INVENTION: Portions of these Genes and Uses Therefor
; FILE REFERENCE: 01997/211003
; CURRENT APPLICATION NUMBER: US/09,888,243
; CURRENT FILING DATE: 2001-06-22
; PRIOR APPLICATION NUMBER: US 09/083,662
; PRIOR FILING DATE: 1998-05-22
; PRIOR APPLICATION NUMBER: US 08/394,189
; PRIOR FILING DATE: 1995-02-24
; PRIOR APPLICATION NUMBER: US 08/282,211
; PRIOR FILING DATE: 1994-07-11
; PRIOR APPLICATION NUMBER: US 07/984,182
; PRIOR FILING DATE: 1992-11-20
; PRIOR APPLICATION NUMBER: US 07/897,788
; PRIOR FILING DATE: 1992-06-12
; NUMBER OF SEQ ID NOS: 30
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 13
; LENGTH: 171
; TYPE: PRT
; ORGANISM: Murine
; FEATURE:
; NAME/KEY: VARIANT
; LOCATION: 117
; OTHER INFORMATION: Xaa - Ala or Val
US-09-888-243-13

Qry Match.   100.0%, Score 24; DB 10; Length 171;
Best Local Similarity 100.0%; Prd. No. 61;
Matches   5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Q7   1 KEELM 5
         |||||
Db   69 KEELM 73

RESULT 7
US-09-888-243-26
; Sequence 26, Application US/09888243
; Patent No. US20020136714A1
; GENERAL INFORMATION:
; APPLICANT: Horvitz, H. Robert
; APPLICANT: Yuan, Junying
; APPLICANT: Shaham, Shai
; TITLE OF INVENTION: Relatedness of Human Interleukin-1beta
; TITLE OF INVENTION: Convertase Gene to a C. Elegans Cell Death Gene, Inhibitory
; TITLE OF INVENTION: Portions of these Genes and Uses Therefor
; FILE REFERENCE: 01997/211003
; CURRENT APPLICATION NUMBER: US/09,888,243
; CURRENT FILING DATE: 2001-06-22
; PRIOR APPLICATION NUMBER: US 09/083,662
; PRIOR FILING DATE: 1998-05-22
; PRIOR APPLICATION NUMBER: US 08/394,189
; PRIOR FILING DATE: 1995-02-24
; PRIOR APPLICATION NUMBER: US 08/282,211
; PRIOR FILING DATE: 1994-07-11
; PRIOR APPLICATION NUMBER: US 07/984,182
; PRIOR FILING DATE: 1992-11-20
; PRIOR APPLICATION NUMBER: US 07/897,788
; PRIOR FILING DATE: 1992-06-12
; NUMBER OF SEQ ID NOS: 30
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 26
; LENGTH: 172
; TYPE: PRT
; ORGANISM: Murine
; FEATURE:
; NAME/KEY: VARIANT
; LOCATION: 118
; OTHER INFORMATION: Xaa - Any Amino Acid
US-09-888-243-26

Qry Match.   100.0%, Score 24; DB 10; Length 172;
Best Local Similarity 100.0%; Prd. No. 61;
Matches   5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy   1 KEELM 5
         |||||
Db   69 KEELM 73

RESULT 8
US-09-888-243-27
; Sequence 27, Application US/09888243
; Patent No. US20020136714A1
; GENERAL INFORMATION:

; APPLICANT: Horvitz, H. Robert
; APPLICANT: Yuan, Junying
; APPLICANT: Shaham, Shai
; TITLE OF INVENTION: Relatedness of Human Interleukin 1beta
; TITLE OF INVENTION: Convertase Gene to a C. Elegans Cell Death Gene, Inhibitory
; TITLE OF INVENTION: Portions of these Genes and Uses Therefor
; FILE REFERENCE: 01997/211003
; CURRENT APPLICATION NUMBER: US/09/888,243
; CURRENT FILING DATE: 2001-06-22
; PRIOR APPLICATION NUMBER: US 09/083,662
; PRIOR FILING DATE: 1998-05-22
; PRIOR APPLICATION NUMBER: US 08/394,189
; PRIOR FILING DATE: 1995-02-24
; PRIOR APPLICATION NUMBER: US 08/282,211
; PRIOR FILING DATE: 1994-07-11
; PRIOR APPLICATION NUMBER: US 07/984,182
; PRIOR FILING DATE: 1992-11-20
; PRIOR APPLICATION NUMBER: US 07/897,788
; PRIOR FILING DATE: 1992-06-12
; NUMBER OF SEQ ID NOS: 30
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 27
;   LENGTH: 172
;   TYPE: PRT
;   ORGANISM: Murine
;   FEATURE:
;   NAME/KEY: VARIANT
;   LOCATION: 118
;   OTHER INFORMATION: Xaa - Any Amino Acid
US-09-888-243-27

Query Match   100.0%; Score: 24; DB 10; Length 172;
Best Local Similarity 100.0%; Pred. No. 61;
Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy   1 KEELM 5
         |||||
Db   69 KEELM 73

RESULT 9
US 10 060-425-15
; Sequence 15, Application US/10060425
; Patent No. US20020164650A1
; GENERAL INFORMATION:
; APPLICANT: Hiebsch, Ronald
; TITLE OF INVENTION: Methods of Assessing Wolframin Protein Activity
; FILE REFERENCE: 00450.US1
; CURRENT APPLICATION NUMBER: US/10/060,425
; CURRENT FILING DATE: 2002-01-30
; PRIOR APPLICATION NUMBER: 60/266,385
; PRIOR FILING DATE: 2001-02-02
; NUMBER OF SEQ ID NOS: 17
; SOFTWARE: PatentIn version 3.1
; SEQ ID No 15
;   LENGTH: 434
;   TYPE: PRT
;   ORGANISM: Alysia
US 10 060-425-15

Query Match   100.0%; Score: 24; DB 9; Length 434;
Best Local Similarity 100.0%; Pred. No. 1.6e+02;
Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy   1 KEELM 5
         |||||
Db 418 KEELM 422

RESULT 10
US-09-827-040-7
; Sequence 7, Application US/09827040
; Patent No. US20010024792A1

; GENERAL INFORMATION:
; APPLICANT: Chen, Hong
; APPLICANT: Meyer, Joanne
; TITLE OF INVENTION: Method of Detecting Risk of Type II Diabetes Based on
; TITLE OF INVENTION: Mutations Found in Carboxypeptidase E
; FILE REFERENCE: 5800-14, 03580/174130
; CURRENT APPLICATION NUMBER: US/09/827,040
; CURRENT FILING DATE: 2001-04-05
; PRIOR APPLICATION NUMBER: 09/233,989
; PRIOR FILING DATE: 1999-01-19
; PRIOR APPLICATION NUMBER: 60/105,102
; PRIOR FILING DATE: 1998-10-21
; NUMBER OF SEQ ID NOS: 10
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 7
;   LENGTH: 434
;   TYPE: PRT
;   ORGANISM: alysia
;   FEATURE:
;   OTHER INFORMATION: carboxypeptidase E
US-09-827-040-7

Query Match   100.0%; Score: 24; DB 10; Length 434;
Best Local Similarity 100.0%; Pred. No. 1.6e+02;
Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy   1 KEELM 5
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Db 418 KEELM 422

RESULT 11
US-09-888-243-28
; Sequence 28, Application US/09888243
; Patent No. US20020136714A1
; GENERAL INFORMATION:
; APPLICANT: Horvitz, H. Robert
; APPLICANT: Yuan, Junying
; APPLICANT: Shaham, Shai
; TITLE OF INVENTION: Relatedness of Human Interleukin-1beta
; TITLE OF INVENTION: Convertase Gene to a C. Elegans Cell Death Gene, Inhibitory
; TITLE OF INVENTION: Portions of these Genes and Uses Therefor
; FILE REFERENCE: 01997/211003
; CURRENT APPLICATION NUMBER: US/09/888,243
; CURRENT FILING DATE: 2001-06-22
; PRIOR APPLICATION NUMBER: US 09/083,662
; PRIOR FILING DATE: 1998-05-22
; PRIOR APPLICATION NUMBER: US 08/394,189
; PRIOR FILING DATE: 1995-02-24
; PRIOR APPLICATION NUMBER: US 08/282,211
; PRIOR FILING DATE: 1994-07-11
; PRIOR APPLICATION NUMBER: US 07/984,182
; PRIOR FILING DATE: 1992-11-20
; PRIOR APPLICATION NUMBER: US 07/897,788
; PRIOR FILING DATE: 1992-06-12
; NUMBER OF SEQ ID NOS: 30
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 28
;   LENGTH: 451
;   TYPE: PRT
;   ORGANISM: Murine
US-09-888-243-28

Query Match   100.0%; Score: 24; DB 10; Length 451;
Best Local Similarity 100.0%; Pred. No. 1.7e+02;
Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy   1 KEELM 5
         |||||
Db 349 KEELM 353

RESULT 12

US-10-060-425-4
; Sequence 4, Application US/10060425
; Patent No. US2002016465OA1
; GENERAL INFORMATION:
;   APPLICANT: Hiebsch, Ronald
;   TITLE OF INVENTION: Methods of Assessing Wolframin Protein Activity
;   FILE REFERENCE: 00450.US1
;   CURRENT APPLICATION NUMBER: US/10/060,425
;   CURRENT FILING DATE: 2002-01-30
;   PRIOR APPLICATION NUMBER: 60/266,385
;   PRIOR FILING DATE: 2001-02-02
;   NUMBER OF SEQ ID NOS: 17
;   SOFTWARE: Patentin version 3.1
; SEQ ID NO 4
;   LENGTH: 476
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-10-060-425-4

Query Match   100.0%; Score 24; DB 9; Length 476;
Best Local Similarity 100.0%; Pred. No. 1.8e+02;
Matches   5; Conservative   0; Mismatches   0; Indels   0; Gaps   0;

Qy   1 KEELM 5
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Db   460 KEELM 464

RESULT 13
US-10-060-425-12
; Sequence 12, Application US/10060425
; Patent No. US2002016465OA1
; GENERAL INFORMATION:
;   APPLICANT: Hiebsch, Ronald
;   TITLE OF INVENTION: Methods of Assessing Wolframin Protein Activity
;   FILE REFERENCE: 00450.US1
;   CURRENT APPLICATION NUMBER: US/10/060,425
;   CURRENT FILING DATE: 2002-01-30
;   PRIOR APPLICATION NUMBER: 60/266,385
;   PRIOR FILING DATE: 2001-02-02
;   NUMBER OF SEQ ID NOS: 17
;   SOFTWARE: Patentin version 3.1
; SEQ ID NO 12
;   LENGTH: 476
;   TYPE: PRT
;   ORGANISM: Rattus sp.
US-10-060-425-12

Query Match   100.0%; Score 24; DB 9; Length 476;
Best Local Similarity 100.0%; Pred. No. 1.8e+02;
Matches   5; Conservative   0; Mismatches   0; Indels   0; Gaps   0;

Qy   1 KEELM 5
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Db   460 KEELM 464

RESULT 14
US-10-060-425-14
; Sequence 14, Application US/10060425
; Patent No. US2002016465OA1
; GENERAL INFORMATION:
;   APPLICANT: Hiebsch, Ronald
;   TITLE OF INVENTION: Methods of Assessing Wolframin Protein Activity
;   FILE REFERENCE: 00450.US1
;   CURRENT APPLICATION NUMBER: US/10/060,425
;   CURRENT FILING DATE: 2002-01-30
;   PRIOR APPLICATION NUMBER: 60/266,385
;   PRIOR FILING DATE: 2001-02-02
;   NUMBER OF SEQ ID NOS: 17
;   SOFTWARE: Patentin version 3.1
; SEQ ID NO 14
;   LENGTH: 476
;   TYPE: PRT
;   ORGANISM: Murinae gen. sp.
US-10-060-425-14

Query Match   100.0%; Score 24; DB 9; Length 476;
Best Local Similarity 100.0%; Pred. No. 1.8e+02;
Matches   5; Conservative   0; Mismatches   0; Indels   0; Gaps   0;

Qy   1 KEELM 5
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Db   460 KEELM 464

RESULT 15
US-09-827-040-2
; Sequence 2, Application US/09827040
; Patent No. US20010024792A1
; GENERAL INFORMATION:
;   APPLICANT: Meyer, Joanne
;   TITLE OF INVENTION: Method of Detecting Risk of Type II Diabetes Based on
;   TITLE OF INVENTION: Mutations Found in Carboxypeptidase E
;   FILE REFERENCE: 5800-14, 03580/174130
;   CURRENT APPLICATION NUMBER: US/09/827,040
;   CURRENT FILING DATE: 2001-04-05
;   PRIOR APPLICATION NUMBER: 09/233,989
;   PRIOR FILING DATE: 1999-01-19
;   PRIOR APPLICATION NUMBER: 60/105,102
;   PRIOR FILING DATE: 1998-10-21
;   NUMBER OF SEQ ID NOS: 10
;   SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 2
;   LENGTH: 476
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-827-040-2

Query Match   100.0%; Score 24; DB 9; Length 476;
Best Local Similarity 100.0%; Pred. No. 1.8e+02;
Matches   5; Conservative   0; Mismatches   0; Indels   0; Gaps   0;

Qy   1 KEELM 5
          |||||
Db   460 KEELM 464

Search completed: January 16, 2003, 17:00:07
Job time : 4.07143 secs